

replacement paragraphs. A marked-up version of these replacement paragraphs is attached on a separate sheet(s).

On page 4, 3rd full paragraph starting at line 9, please delete and replace the current version of the paragraph, with the following replacement paragraph:

This object is achieved by the provision of a DNA sequence which exhibits at least a 70% homology to the sequence (SEQ ID NO: 13) as shown in Fig. 12, and which codes for a polypeptide having the biological activity of the enzyme amorphadiene synthase.

On page 9, 9th full paragraph starting with “Fig. 8”, please delete and replace the current version of the paragraph, with the following replacement paragraph:

Fig. 8: Nucleotide sequence (SEQ ID NO: 9) and deduced amino acid sequence (SEQ ID NO: 10) of the probe (538 bp) generated by PCR with primers A and B.

On page 9, 11th full paragraph starting with “Fig. 10”, please delete and replace the current version of the paragraph, with the following replacement paragraph. Please note that both the marked-up version and the clean version contain underlining that is to be retained and does not constitute an amendment.

Fig. 10: Nucleotide sequence (SEQ ID NO: 11) and deduced amino acid sequence (SEQ ID NO: 12) of a positive clone (amorphadiene synthase encoding gene) isolated from the cDNA library of induced A.annua. The sequence (SEQ ID NO: 11) is flanked with EcoRI (NotI) adapters (Gibco BRL).

On page 10, starting with “Fig. 12”, please delete and replace the current version of the paragraph, with the following replacement paragraph. Please note that both the marked-up version and the clean version contain underlining that is to be retained and does not constitute an amendment.

Fig. 12: Nucleotide sequence (SEQ ID NO: 13) and deduced amino acid sequence (SEQ ID NO: 14) of the amorphadiene synthase encoding gene, between start and stop codon (flanked by NcoI and BamHI sites, respectively), obtained by PCR with primers C and D.

On page 17, starting at line 1, please delete and replace the current version of the paragraph, with the following replacement paragraph.

5'-pGTCGACGCGGCCGCG-3' (SEQ ID NO: 1)

On page 17, starting at line 2, please delete and replace the current version of the paragraph, with the following replacement paragraph.

3'-CAGCTGCGCCGGCGCTTAA-OH-5' (SEQ ID NO: 2)

On page 17, starting at line 33 and bridging page 18 and ending at line 9, please delete and replace the current version of the paragraph, with the following replacement paragraph. Please note that both the marked-up version and the clean version contain underlining that is to be retained and does not constitute an amendment.

For functional expression the cDNA clone was subcloned in frame into the expression vector pET lld (Stratagene). To introduce suitable restriction sites for subcloning, the gene was amplified by PCR using a sense primer (primer C)

5'-GTCGACAAACCATGGCACTTACAGAA G-3' (SEQ ID NO: 3) (introducing a NcoI site at the start codon **ATG**) and an anti-sense primer (primer D):

5'-GGATGGATCCTCATATACTCATAGGATAAACG-3' (SEQ ID NO: 4) (introducing a BamHI site directly behind the stop codon **TGA**). The PCR reaction was performed under standard conditions. After digestion with BamHI and NcoI, the PCR product (Fig. 12) and the expression vector pET 11d were gel purified and ligated together to yield a construct as revealed in Fig. 11.

On page 20, 2nd full paragraph starting at line 9, please delete and replace the current version of the paragraph, with the following replacement paragraph. Please note that both the marked-up version and the clean version contain underlining that is to be retained and does not constitute an amendment.

Analogous to EXAMPLE 3, suitable restriction sites for subcloning were introduced by using PCR with a sense primer (primer G) 5'-GA GGA TCC ATG TCA CTT ACA GAA-3' (SEQ ID NO: 5) (introducing a BamHI site preceding the start codon **ATG**) and an anti-sense primer (primer H) 5'-AT GGA TCC TCA TAT ACT CAT AGG A-3' (SEQ ID NO: 6) (introducing a BamHI site directly behind the stop codon **TGA**). After digestion with BamHI the PCR product and the plant-expression cassette pLV399 were gel purified and ligated to provide the gene encoding amorpho-4, 11-diene synthase with the cauliflower mosaic virus 35S promoter and a nopaline synthase transcription terminator. The plant-expression cassette pLV399 is a pUC 19 vector (Yanisch-Perron, C. et al., Gene 33, 103-119 (1985)) in which the multiple cloning site (polylinker) is replaced by a CaMV 35 S promoter BamHI fused to a nos-tail (terminator) flanked by the 'unique' sites; EcoRI, KpnI, XhoI, and a HindIII site downstream from the promoter and EcoRI, XhoI, PstI, SphI, KpnI, HindIII upstream from the terminator. The orientation of the amorpho-4, 11-diene encoding

gene in pLV399 was checked by restriction analysis with PstI and NdeI. After partial digestion of this construct with KpnI the amorpho-4, 11-diene encoding gene flanked by the 35S promotor and nos terminator was ligated into the KpnI digested binary vector pCGN1548.

On page 22, starting at line 29 and bridging page 23 and ending at line 3, please delete and replace the current version of the paragraph, with the following replacement paragraph. Please note that both the marked-up version and the clean version contain underlining that is to be retained and does not constitute an amendment.

For functional expression the cDNA clone was subcloned into the inducible expression vector pYES2 (episomal vector, Invitrogen) and the constitutive expression vector (integrating the gene construct into the genome) pGAPZ A (Invitrogen). To introduce suitable restriction sites for subcloning, the gene was amplified by PCR using a sense primer (primer E) 5'-CGA GAA TTC ATG TCA CTT ACA G-3' (**SEQ ID NO: 7**) (introducing a ExoRI site preceding the start codon **ATG**) and an anti-sense primer (primer F) 5'-GGAT CTC GAG TCA TAT ACT CAT-3' (**SEQ ID NO: 8**) (introducing a BamHI site directly behind the stop codon **TGA**). Subcloning of the PCR product into pYES2 and pGAPZ A was done in a way analogue to Example 3.

IN THE DRAWINGS:

Figs. 8, 10 and 12 have been amended to include the addition of SEQ ID NOs which are indicated in red. Copies of Figs. 8, 10 and 12 are attached on separate sheets. The correction of Figs. 8, 10 and 12 will be made formally when formal figures are filed at a later date.